## Genospectra QuantiGene Study Plan and SOP For MAQC

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# Genospectra QuantiGene Platform

### Test Site

Genospectra, Inc.

Contact person: Yuling Luo

Address: 6519 Dumbarton Circle, Fremont, CA 94555

Phone: 510.818.2623

Email: yluo@genospectra.com

### Materials Required

### Genospectra Products

- QuantiGene kits
- Gene-specific probe sets for over 200 genes

### Non-Genospectra Products

- Total RNA, Brain (Ambion)(AB)
- Total RNA, Universal Human Reference RNA (Stratagene) (UHRR)
- Total RNA, Mixture of 75% AB + 25% UHRR
- Total RNA, Mixture of 25% AB + 75% UHRR

#### Equipment and consumables

- General laboratory equipment include a 96-well plate-based luminometer (Lmax, Molecular Device) and a hybridization incubator (Hybaid) capable of maintaining temperature between 46°C and 53°C.
- General consumables are listed in the QuantiGene assay SOP manual

#### User Manuals

• Genospectra QuantiGene Assay SOP Manual

### Experimental Design

Minimal of 200 genes will be selected for the QuantiGene assay and agreed to by MAQC. The 200 gene will be matched within the ABI's 1,000 gene list, which will be matched within the list of 5,000 RefSeq genes shared by all microarray platforms. The 200 genes will be selected to cover different expression level, fold changes, and some genes showing cross-platform discrepancy in the pilot study.

Agreement need to be reached among Genospectra, ABI, and Stratagene for the platform validation assays. Three, four or five replicate measurements for each gene can be processed for each of the following mRNA samples, depending on the agreement among Genospectra, ABI, and Stratagene:

- 1. UHRR (100%)
- 2. AB (25%) vs. UHRR (75%)
- 3. AB (75%) vs. UHRR (25%)

### Total RNA Preparation and Analysis

It is assumed that all four total RNA samples will be provided in aqueous solution ready for use. The total RNA samples will be stored at  $-80~^{\circ}\text{C}$  and will be used directly in the QuantiGene assay, unless there is a need to dilute the sample further. No additional manipulation is required for the QuantiGene assay. As a result, we rely on the quality assessment of the original total RNA samples by Ambion and Stratagene.

#### QuantiGene® assay conditions:

- 1. Stock total RNA samples will be diluted to  $[40\,\mathrm{ng/ul}]$  in RNase Free water.
  - 2. 100 to 800ng of total RNA will be added to each well of QuantiGene assay, depending on the gene expression level.
- 3. Hybridization assay will be carried out per QuantiGene manual.
  - 4. Background signal for each gene will be assessed in a QuantiGene assay with no sample RNA added, just the probe set.

### Data Acquisition and Analysis

QuantiGene assay signal is measured by a luminometer, and the Relative Luminescent Unit (RLU) data will be directly read from the luminometer machine. The data will be exported into an Excel file and reported directly to MAQC.

For those genes (e.g. selected genes from MAQC pilot II study) whose expression in AB and UHRR is substantially different (e.g. over 100: 1 ratio), we expect an assay signal ratio of 100: 75:25:0 ratio for the four total RNA samples. The closeness to the expected 100:75:25:0 ratio will be a measurement of the assay platform accuracy.

Once the assay platform accuracy is established, the relative ratio of the 200 genes among different samples will be assessed, and their correlation with the data from other platforms will be compared.

## **Overview of the QuantiGene Workflow**

# Day One:

- A. Prepare the QuantiGene® Reagents and gene-specific Probe Set
- B. Prepare RNA samples for the selected genes.
- C. Follow Experimental Layout on Plate Map.

## Day Two:

- D. Prepare the QuantiGene Reagents used for the target mRNA.
- E. Amplify the Signal and Detect target mRNA.
- F. Analyze the Data.

## **QuantiGene Assay SOP**

### **PURPOSE:**

This protocol describes the procedure for performing, single-plex QuantiGene assays with purified RNA using the LMAX luminometer.

### SCOPE:

This procedure applies to all R&D personnel performing a QuantiGene single-plex assay using purified RNA or IVTs as the sample source.

### **KEY SPONSOR AND RESPONSIBILITY:**

The R&D Manager is responsible for the content, updating and implementation of this document.

### **EQUIPMENT:**

- 1. Pipettes, single and multichannel
- 2. Vortex mixer
- 3. LMAX luminometer
- 4. Incubators: 46° +/-1.0°C and 53° C +/- 1.0° C
- 5. Quick spin bench top micro-centrifuge

### MATERIALS

- 1. QuantiGene Assay Kit (at 4°C capture plates, amplifier, label probe, substrate; at 15-30°C amplifier/label probe diluent, lysis mixture)
- 2. Plastic pipette tips, aerosol resistant
- 3. Disposable reagent reservoirs, 25 and 100 mL capacity
- 4. Plate Sealers (Bioexpress, Cat. No. T-3025-8B)
- 5. 10X or 1X Wash Buffer
- 6. 10% Lithium Lauryl Sulfate solution
- 7. Capture Buffer
- 8. 1 mg/mL tRNA (Blocking Agent)
- 9. Purified total RNA, mRNA, or IVT RNA
- 10. Appropriate Probe Set(s) with pooled Label Extenders, pooled Capture Extenders and pooled Blocking Probes
- 11. Nuclease Free water

## **SOP Procedure:**

# Day One:

Step	Action
	Bring the Capture Plate and target-specific probes to room temperature.
1	
2	Check to be sure the <b>Lysis Mixture</b> does not have precipitates. If you observe precipitates, then perform the following: a. Place the bottle in a 37°C water bath. b. Gently swirl contents occasionally until all precipitates dissolve.
3	Prepare the <b>Concentrated Probe Sets</b> for the target RNA of interest by adding 50 µl each of pooled 1X CE, (Capture Extender, 50 fmol/µl), 1X LE, (Labeled Extender, 200 fmol/µl), and 1X BL, (Blocking Probe, 100 fmol/µl), probes to 850 µl of <b>Lysis Mixture</b> (from QuantiGene kit). <b>Note: This preparation of concentrated probe set is enough for one 96-well plate.</b>
4	Prepare Lysis Working Reagent: Add 15 µl of yeast tRNA, to 6 ml of Lysis Mixture
5	Depending upon the expected abundance of the specific mRNA to be measured, dilute the RNA sample with <b>Lysis Working Reagent</b> (see Step 4, above) to a final concentration of 0.02 to 2 $\mu$ g/10 $\mu$ l, if using mRNA dilute to a final concentration of 0.4 to 40 ng/10 $\mu$ l. <b>Note:</b> For a gene with a moderate level of expression, 0.2 $\mu$ g/10 $\mu$ l or 4 ng/10 $\mu$ l of mRNA are recommended.
6	Dispense 50 µl of Capture Buffer, (Catalog # QGR-004-002), to the Capture Plate well
7	Dispense 40 μl of <b>Lysis Mixture</b> to the <b>Capture Plate</b> well.
8	Dispense 10 μl of <b>RNA sample</b> to the <b>Capture Plate</b> well.
9	Dispense 10 µl of Concentrated Probe Sets to the Capture Plate well.
10	Seal the <b>Capture Plate</b> with a plate sealer. Incubation the sealed plate at 53°C for 16-20 hours.

## Day Two:

### **Procedural Guidelines**

- ♦ We recommend preparing the **Substrate**, **Amplifier** and **Label Probe** immediately before use. Prior to use, bring these reagents to room temperature.
- Vortex and briefly centrifuge the Amplifier and Label Probe to bring the contents to the bottom of the tube before use.
- ◆ Calculations below for the Wash Buffer include a 20% overage. Calculations for the Amplifier, Label Probe and Substrate Working Reagents include a 10% overage. Determine the specific volume requirements for your system.

### D. Prepare the Reagents

D. FI	epare the Reagents				
Step	Action				
1	a. b.	Prepare the Wash Buffer:  a. Combine the reagent volumes appropriate for the number of plates used in the assay.  b. Stir gently to mix.  c. Keep mixture at room temperature.			
		Capture	10X bDNA	Distilled	Total
		Plates	Wash Buffer	H <sub>2</sub> 0	Volume
		1	40 ml	360 ml	400 ml
		2	80 ml	720 ml	800 ml
	a.		•		ber of plates used in the
2		Combine the reagen assay. Stir gently to mix. Keep mixture at roor	nt volumes appropi	iate for the num	·
2	a. b.	Combine the reagen assay. Stir gently to mix. Keep mixture at roor  Capture Plates	n temperature.  Amplif	iate for the num	Amplifier
2	a. b.	Combine the reagen assay. Stir gently to mix. Keep mixture at roor	m temperature.  Amplif	iate for the num	·
2	a. b. c.	Combine the reagen assay. Stir gently to mix. Keep mixture at roor  Capture Plates  1 2  repare the Label Pro  Combine the reagen assay. Stir gently to mix.	m temperature.  Amplif 17 22  Dbe Working Re at volumes appropri	ier/Diluent  .6 ml 2.0 ml  agent:	Amplifier 11.6 µl
	a. b. c.	Combine the reagen assay. Stir gently to mix. Keep mixture at roor  Capture Plates  1 2  repare the Label Pro Combine the reagen assay. Stir gently to mix. Keep mixture at roor	Amplif  Amplif  22  Dbe Working Re  It volumes appropri	ier/Diluent 1.6 ml 2.0 ml agent: riate for the num	Amplifier 11.6 μl 22.0 μl ber of plates used in the
	a. b. c.	Combine the reagent assay. Stir gently to mix. Keep mixture at roor  Capture Plates  1 2  repare the Label Processay. Stir gently to mix. Keep mixture at roor  Capture Plates	m temperature.  Amplif 12  Dibe Working Re at volumes appropri	ier/Diluent  1.6 ml  2.0 ml  agent: iate for the num	Amplifier 11.6 μl 22.0 μl ber of plates used in the
	a. b. c.	Combine the reagen assay. Stir gently to mix. Keep mixture at roor  Capture Plates  1 2  repare the Label Pro Combine the reagen assay. Stir gently to mix. Keep mixture at roor	m temperature.  Amplif 17 22 Dbe Working Re at volumes appropri	ier/Diluent 1.6 ml 2.0 ml agent: riate for the num	Amplifier 11.6 μl 22.0 μl ber of plates used in the

**Prepare the Reagents (cont.)** 

٨٠	tion		
	combine the reagent volumes assay.  Stir gently to mix. It is not turbid after the addition of	lumes appropriate for the normal for the Substrate Working the 10% Lithium Lauryl S	orking Reagent to become ulfate.
	Capture Plates	Substrate	10% Lithium Lauryl Sulfate
	1	11.6 ml	34.8 µl
	2	22.0 ml	66.0 µl
	Pre a. b.	<ul> <li>a. Combine the reagent vo assay.</li> <li>b. Stir gently to mix. It is not turbid after the addition of c. Cover the solution with all Capture Plates</li> </ul>	Prepare the Substrate Working Reagent:  a. Combine the reagent volumes appropriate for the massay.  b. Stir gently to mix. It is normal for the Substrate We turbid after the addition of the 10% Lithium Lauryl Sc. Cover the solution with aluminum foil and keep at roo  Capture Plates Substrate  1 11.6 ml

E. Amplify and Detect the Target mRNA

Step	Action	
	Centrifugation Method	r and remove the seal from the plate. s then follow one of the methods below.  Aspiration
	[Highly recommended]	Method
1	<ul> <li>a. Decant the Wash Buffer and Cell Lysate to a sink.</li> <li>b. Wash each well 3 times with</li> </ul>	<ul> <li>a. Aspirate the Wash Buffer and Cell Lysate.</li> <li>b. Wash each well 3 times with 350 µl of Wash Buffer.</li> </ul>
'	350 µl of <b>Wash Buffer.</b> c. Decant the buffer to a sink and keep plate face down on paper towels.	c. Completely remove <b>Wash Buffer</b> from the wells; the aspiration nozzle may be lowered down the side of
	d. Following third wash use centrifugation to remove	the well until it contacts the bottom.  Notes:
	traces of the <b>Wash Buffer</b> from the plate.	Try not to touch the bottom of the well.
	Lay the plate face down on 2 to 3 paper towels and put in the centrifuge bucket. Spin at 240 g for 1 minute.	2. Try to touch the same corner of the well with the pipette tip every time.

c. R	·	and remove the seal from the plate sealer then follow one of the methods below.
	Spin Approach [Highly recommended]	Aspiration Approach
e. f. g. h.	Decant the Wash Buffer and Cell Lysate to a sink. Wash each well 3 times with 350 µl of Wash Buffer. Decant the buffer to a sink and keep plate face down on paper towels. Following third wash use spinning to remove traces of the Wash Buffer from the plate. Lay the plate face down on 2 to 3 paper towels and put in the centrifuge bucket. Spin at 240 g for 1 minute.	<ul> <li>d. Aspirate the Wash Buffer and Cell Lysate.</li> <li>e. Wash each well 3 times with 350 μl of Wash Buffer.</li> <li>f. Completely remove Wash Buffer from the wells; the aspiration nozzle may be lowered down the side of the well until it contacts the bottom.</li> <li>Notes:</li> <li>3. Try not to touch the bottom of the well.</li> <li>4. Try to touch the same corner of the well with the pipette tip every time.</li> </ul>

**Amplify and Detect the Target mRNA (cont.)** 

Step	Action		
2	Immediately add 100 µl of <b>Amplifier Working Reagent</b> to each well.		
3	Reseal and incubate the plate at 46° to 53°C for 60 minutes.		
4	Wash the Capture Plate:  a. Remove the plate from the incubator and remove the seal.  b. Immediately aspirate or decant the Amplifier Working Reagent from the wells and wash each well three times with 350 µl of Wash Buffer.  c. Completely remove Wash Buffer from the wells. (Refer to Step 1.)		
4	Wash the Capture Plate:  d. Remove the plate from the incubator.  e. Immediately aspirate the Amplifier Working Reagent from the wells and wash each well three times with 350 µl of Wash Buffer.  f. Completely remove Wash Buffer from the wells. (Refer to Step 1.)		

5	Immediately add 100 µl of Label Probe Working Reagent to each well.		
6	Reseal and incubate the plate at 46° to 53°C for 60 minutes.		
	Wash the Capture Plate:		
	a. Remove the plate from the incubator and remove the seal.		
7	b. Immediately aspirate or decant the <b>Label Probe Working Reagent</b> from the wells and wash each well three times with 350 µl of <b>Wash Buffer</b> .		
	c. Completely remove <b>Wash Buffer</b> from the wells. (Refer to Step 1.)		
	Note: After hybridizing with Label Probe do not let the plate dry for more than 5 minutes to avoid partial inactivation of the AP activity.		
	Wash the Capture Plate:		
	d. Remove the plate from the incubator.		
7	e. Immediately aspirate the <b>Label Probe Working Reagent</b> from the wells and wash each well three times with 350 µl of <b>Wash Buffer</b> .		
	f. Completely remove <b>Wash Buffer</b> from the wells. (Refer to Step 1.)		
	Note: After hybridizing with Label Probe do not let the plate dry for more than 5 minutes to avoid partial inactivation of the AP activity.		
8	Immediately add 100 µl of <b>Substrate Working Reagent</b> to each well.		
9	Reseal and incubate at 46° to 53°C for 30 minutes.		
10	Remove the plate from the incubator and transfer the plate to a luminometer (without a plate heater) and allow the plate to cool for 5 minutes. <b>Alternatively</b> , for a luminometer with a plate heater, allow the plate to equilibrate in the tray for 5 minutes.		
11	Measure the luminescence of each well.		

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